

**Diet composition has a greater effect on larval and juvenile polar cod (*Boreogadus saida*)
growth rates than temperature in a warming Pacific Arctic**

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Keywords: *Boreogadus saida*, Chukchi Sea, Diet, Temperature, Growth, Otolith, Compound-specific stable isotope analysis of amino acids

Abstract

The relative influences of temperature and diet on growth of marine fish are especially relevant in high-latitude ecosystems where warming temperatures and loss of sea ice alter the thermal and feeding environment for cold-adapted species. Larval and juvenile fish may be particularly vulnerable to warming since access to favorable habitats in the spring and summer is necessary for growth and overwinter survival. Therefore, we focused on three warm, low-ice years (2017-2019) in the Chukchi Sea and on a nodal forage fish species in Arctic food webs (polar cod: *Boreogadus saida*) to better understand the influences of warming temperatures and diet on growth during the first year of life. Paired diet data from gut content analysis and daily otolith-derived growth rates indicated that diet had a greater impact on growth than temperature. Generalized additive models that included diet information explained a large portion of variability in recent growth [5-d average preceding capture; coefficient of determination (R^2) 0.43-0.45] compared to those that excluded diet but incorporated temperature and salinity ($R^2=0.06-0.07$) and a year effect ($R^2=0.20-0.30$). Relationships between diet and growth suggest that polar cod fed on a range of taxa during warm years but benefited from access to remnant sea-ice associated prey such as *Calanus glacialis* copepods. Larval and juvenile polar cod may be resilient to warming temperatures if preferential high-quality prey persists into the summer, but sea-ice retreat and inundation of subarctic zooplankton earlier in the year could have detrimental impacts on polar cod growth and survival.

1. Introduction

Fast growth and large body size are often associated with increased survival of marine fish during the first year of life due to lower risk of predation and presumed higher body condition (Heintz and Vollenweider, 2010; Robert et al., 2023; Sogard, 1997). Growth variation during the early life stages is largely driven by a combination of temperature (Atkinson, 1994) and food availability to meet metabolic demands (Mazur et al., 2007; Sreenivasan and Heintz, 2016). Under this paradigm, environmental change or small-scale variation in local conditions may have direct or indirect effects on growth and survival.

High-latitude ecosystems are undergoing rapid environmental change where warming water temperatures and changes in primary and secondary productivity affect the thermal and

feeding environment for the early life stages of marine fish (Kimmel et al., 2023; Leu et al., 2011; Søreide et al., 2010). In the Arctic, earlier summer sea-ice retreat alters the composition of basal resources from largely sea-ice algae to a greater contribution from subarctic primary producers (Lee et al., 2019). Loss of sea ice also leads to lower abundance of large lipid-rich copepods such as *Calanus glacialis* that are important prey for larval and juvenile fish (Bouchard and Fortier, 2020; Spear et al., 2020; Strasburger et al., 2014). For polar cod [*Boreogadus saida* (Lepechin, 1774); Gadidae; alternatively referred to as “Arctic cod”], the most abundant forage fish in the Arctic (Geoffroy et al., 2016; Kono et al., 2016; Welch et al., 1992), warmer temperatures and decreases in *C. glacialis* could affect growth by exposing fish to temperatures above their thermal optimum (Laurel et al., 2016) and limiting preferred prey (Bouchard and Fortier, 2020).

Polar cod serve as a link between sea-ice algal primary producers and higher trophic levels such as seals, whales, and sea birds (Hop and Gjøsæter, 2013; Kohlbach et al., 2017), which merits a better understanding of the impacts of warming and diet shifts on growth and survival of this nodal species (Mueter et al., 2016; Welch et al., 1992). Polar cod life history is closely tied to sympagic ecosystems such that large-scale loss of sea-ice habitats may lead to overall declines in abundance and biomass (Geoffrey et al., 2023) or restrict polar cod distributions due to borealization of their southern range margin (Axler et al., 2023; Levine et al., 2023; Randall et al., 2019; Spear et al., 2020). Laboratory studies highlight the potential nuanced interaction between warming and growth due to metabolic constraints, whereby polar cod larval growth rates increase with temperature up to ~5°C-7°C, although mortality also increases (Koenker et al., 2018). In the natural environment, extensive declines in sea ice and warming temperatures will likely increase metabolic demands for larval and juvenile polar cod (David et al., 2022) while simultaneously reducing access to favorable large, lipid rich copepod prey that rely on sea-ice algal productivity (Bouchard and Fortier, 2020; Søreide et al., 2010). There is evidence to suggest benefits of early ice break-up and early hatch to extend the growing season (Bouchard et al., 2017), but warmer temperatures are generally associated with reduced body size and condition of juveniles, with consequences to pre-winter provisioning and overwinter survival (Copeman et al., 2017; 2022b). Despite overarching detrimental effects of loss of sea ice on polar cod, variation in spatial distributions, diet resources, hatch timing, and growth suggest

that the species may be adaptable, or have complex responses, to warming (Bouchard and Fortier, 2011; Bouchard et al., 2017; Goldstein et al., 2023; Levine et al., 2023).

Few empirical studies have addressed the combined influences of temperature and diet on polar cod growth and size during the early life stages, particularly in the Chukchi Sea that serves as a Pacific “gateway” to the Arctic (David et al., 2022; Geoffroy et al., 2023). There are even fewer studies in the Chukchi Sea continental shelf ecosystem addressing the influence of environmental conditions on individual larval and juvenile polar cod growth (Chapman et al., 2023) despite the prevalence of age-0 polar cod on the productive shelf, and their important trophic role in Arctic food webs (Levine et al., 2023). To help fill this gap in understanding about polar cod vulnerabilities to a changing Arctic, we use otolith-derived growth rates and morphological diet data from larval and juvenile fish collected in the Chukchi Sea. We focus on three study years (2017-2019) that provide possible representations of a “future” Arctic with reduced sea ice, warm temperatures, and an inundation of subarctic water masses (Goldstein et al., 2023; Levine et al., 2023; Spear et al., 2020) to determine the influences of temperature and diet composition on growth and body size of polar cod during the early life stages. Warm temperatures combined with reduced access to high quality prey could lead to predominantly negative relationships between temperature and growth for cold-adapted polar cod. We hypothesize that during the three warm study years, polar growth rates will be suppressed at the highest observed temperatures, and that potential positive relationships between temperature and growth at moderately warm temperatures (Koenker et al., 2018) will be mediated by access to ice-associated high-quality prey.

2. Methods

2.1 Survey data and specimen collection

Oceanographic data and polar cod samples were collected aboard surveys conducted in the late summer months (August-September) of 2017, 2018, and 2019 (Supplemental Table 1) in the Chukchi Sea (Figure 1). Vertical profiles of temperature and salinity were measured using Sea-Bird Electronics (SBE) 911plus conductivity, temperature, depth recorders (CTD) just prior to or immediately after net tows or a SBE 49 FastCAT CTD affixed to ichthyoplankton nets. CTD data were processed using SBE software with values averaged in 1 m depth bins. Fish were

collected using paired 60-cm (505 µm mesh) and 20-cm (153 µm mesh) bongo nets that were towed obliquely from the surface to 10 m off the bottom or a maximum depth of 200 m (Matarese et al., 2003). All fish samples were sorted from one side of each paired net to preserve the other side for quantitative catch data. To increase sample sizes in 2018, larval and juvenile specimens [$n=3$; < 35 mm standard length; (Vestfals et al., 2021)] were opportunistically collected using a Methot frame trawl (2.23 m by 2.23 m square-frame opening, 2 mm hexagonal mesh, and 500 µm codend mesh; Methot, 1986). The Methot trawl was towed for 8 to 21 minutes, targeting the pelagic scattering layer or the chlorophyll maximum. Samples were stored at -80°C . Thawed fish were measured in the laboratory (standard length, mm). Stomachs were removed and placed in 70% ethanol to prevent degradation.

2.2 Otolith-derived age and growth data

Lapillar and sagittal otoliths were extracted for daily age estimation since lapillus daily increment formation has been confirmed for polar cod (Bouchard and Fortier, 2011) but sagittae are often used for daily age estimation of gadids due to interpretability (Almeida et al., 2024; Dougherty, 2008). Lapilli were mounted in Crystalbond™ 509 thermoplastic cement with no other preparation. Sagittae were transverse thin sectioned by mounting otoliths in Crystalbond™ 509 and manually polishing with fine grit lapping film (ÅngströmLap®; 3 µm) followed by a buffing cloth and 0.05 µm Buehler MasterPrep polishing suspension. Lapilli and sagittae were aged using a compound microscope and a 100× oil immersion objective. Otoliths were aged blindly (no knowledge of fish length or prior age estimates) by a single reader by imaging otoliths (focused on the primordium) and annotating otolith daily increments using ImagePro 10 software (Media Cybernetics). A new image was obtained for each age estimate, with a minimum of two age estimates for each fish. A specimen was retained for analyses if the coefficient of variation among all age estimates was $<10\%$. Final ages were averaged and rounded to integer age. Otoliths were aged from a total of 83 fish, and 72 specimens met the agreement criteria for inclusion in analyses (Supplemental Table 1). Comparisons of fish daily age estimates obtained from sagittae and lapilli ($n=68$ fish) showed good agreement, supporting the use of either otolith for age determination [ordinary least squares regression (OLS): $\text{Age}_{\text{lapillus}} \sim 0.97 * \text{Age}_{\text{sagitta}} + 1.86$; $R^2=0.97$, $p < 0.001$].

The assumption that daily otolith increment widths are correlated with somatic growth rates (Campana 1992; Sponaugle 2009) of polar cod was supported by a correlation between lapilli radii [longest growth axis (posterior)] and fish lengths measured from a random subset of specimens ($n=47$; OLS: otolith radius = $1.32 * \text{daily age} + 12.58$; $R^2=0.82$, $p < 0.001$). Lapillus increment widths were measured along the posterior growth axis to estimate growth rates from three time periods during the early life history of each fish: (1) Recent growth was defined as daily increment widths averaged across the most recent 5 full days of growth prior to capture (excludes the partial increment on the day of capture; $\mu\text{m/d}$). (2) Past growth was defined as average daily increment widths on the preceding 6-10 days before capture ($\mu\text{m/d}$). (3) Early growth was measured as the lapillus otolith radius from the primordium to day 10 post-hatch (e.g., $\mu\text{m}/10\text{d}$; images were focused on the primordium). Growth preceding capture (recent and past growth) were estimated from lapillus images focused on the edge of each otolith to ensure a comparable focal plane among all specimens. For these measurements, a transect was drawn from the otolith edge to the primordium along the growth axis and daily increment widths were measured (starting from the otolith edge) until increments were difficult to discern due to the focal plane. This resulted in differences in the total number of days before capture measured for each specimen. However, all specimens but two included measurements for the most recent 10 days preceding capture.

2.3 Diet data and analyses

Morphological diet data were obtained by removing the alimentary canal from each fish and using minuten pins to separate zooplankton prey in glycerol (Llopiz and Cowen, 2009). Prey were identified to the lowest taxonomic level possible and enumerated using a stereomicroscope with a camera and ImagePro 10 software. Copepodite and adult copepods were not differentiated due to digestion, and some prey taxa were grouped at a higher taxonomic level for simplification of analysis. Diet data (raw data as numerical counts of each taxonomic group) were visualized and simplified with Principal Components Analysis (PCA) using taxa whose counts comprised at least 1% of the total observations among all taxa. A Hellinger transformation (square root of relative abundance) was conducted on the diet count data prior to the PCA to address spurious similarities among specimens due to shared absence of species in diet data (double-zero problem; Legendre and Gallagher, 2001; Legendre and Legendre, 2012).

To better link changes in diet with potential declines in sea ice and warming temperatures, we extended work presented in Goldstein et al. (2023; see Methods section for detailed description of methods and analyses) to compare gut content diet data with available $\delta^{13}\text{C}$ compound-specific stable isotope analysis of amino acid (CSIA-AA) data from homogenized polar cod tissues (excluding heads and stomachs) for a subset of fish from 2017 and 2018. The goal of this analysis was to better understand whether changes in diet were associated with changes in food web basal carbon resources such as sea-ice-algal derived carbon or pelagic phytoplankton. Broadly, CSIA-AA values from essential amino acids exhibit minimal or no trophic fractionation such that CSIA-AA fingerprints (row-centered CSIA-AA for each specimen or observation) mirror CSIA-AA fingerprints from carbon sources (Larsen et al., 2013). Potential carbon sources considered for this analysis were sea-ice algae (iPOM; determined from ice-associated particulate organic matter), pelagic primary producers (pPOM; determined from pelagic particulate organic matter), microalgae, *Synechococcus* spp. and dinoflagellates [typically subarctic taxa; (Flombaum et al., 2013)], and bacteria. We present a Linear Discriminant Analysis adapted from Goldstein et al. (2023) to visualize previously established relationships between CSIA-AA fingerprints from carbon sources and polar cod samples. Additional details that are specific to this study are available in Goldstein et al. (2023). We then used a multivariate redundancy analysis (RDA) with Euclidean distance to determine whether changes in diet corresponded with changes in basal carbon resources for polar cod. RDA is a constrained ordination method where ordination vectors of response variables are linear combinations of the predictor variables (Legendre and Legendre, 2012). Here, we incorporated PCA axis 1, PCA axis 2, and PCA axis 3 scores from gut content analyses as predictor variables. Response variables were row-centered CSIA-AA fingerprints for Isoleucine (Ile), Leucine (Leu), and Valine (Val) from polar cod samples. Other amino acids were excluded from the analyses due to multicollinearity (variance inflation factors: VIFs >3).

2.4 Influence of environment and diet on otolith-derived growth

Analysis of Variance (ANOVA) followed by Tukey's Honestly Significant Difference (Tukey) post-hoc tests were used to determine whether there were average differences in early growth, past growth, recent growth, and diet PCA axis 1 scores (since axis 1 encompasses the most variability in the data) among years. ANOVA tests using a significance threshold of $p <$

0.05 were conducted after confirming that growth metrics were not correlated with age and confirmation that data followed a normal distribution and exhibited homogeneity of variance (Sokal and Rohlf, 1995).

Generalized additive models (GAMs; Gaussian distribution and “identity” link function) were then used to test for potential drivers of variation in growth. Specifically, GAMs focused on the influence of intrinsic (i.e., age and early growth) and extrinsic (i.e., temperature, salinity, diet) factors on growth rates. Since diet and *in situ* data only represent recent conditions, the focal GAM response variable was recent growth (5 days preceding to capture). A second set of models was also tested with past growth (6-10 days preceding to capture) to determine if conditions at capture were correlated with longer-term patterns in growth. For each response variable, two sets of models were tested to focus on the influences of intrinsic factors, environmental conditions, and diet on growth (Equation 1) or the influences of only intrinsic factors and environmental conditions on growth (Equation 2).

$$\text{Equation 1: Growth} = s(S_{200}) + s(T_{20}) + s(S_{20}) + s(\text{Age}) + s(\text{Growth}_{\text{early}}) + s(\text{PC1}_{\text{diet}}) + s(\text{PC2}_{\text{diet}}) + s(\text{PC3}_{\text{diet}})$$

$$\text{Equation 2: Growth} = s(T_{200}) + s(S_{200}) + s(T_{20}) + s(S_{20}) + s(\text{Age}) + s(\text{Growth}_{\text{early}})$$

Here, $s()$ refers to smooth terms with a thin plate spline basis and knots set to 4 to avoid overfitting. The response variable, Growth, is recent growth or past growth determined from otolith increment widths ($\mu\text{m}/\text{day}$). T_{200} and S_{200} are temperature ($^{\circ}\text{C}$) and salinity (PSU), respectively, averaged over the top 200 m of the water column which represented the entire water column at many of the stations (Fig. 1) and encompasses the entire depth range of the bongo net tows where fish were collected. T_{20} and S_{20} are surface temperature and salinity, respectively, averaged over the top 20 m of the water column. The Chukchi Sea water column in the summer months is often stratified with a typical surface mixed layer depth of ~ 20 m (Weingartner et al., 2013). Water mass composition evolves throughout the season (Lin et al., 2019). However, surface waters are generally comprised of cool low salinity sea-ice melt water and warmer Alaskan coastal (Pacific-origin) water masses, whereas deeper depths are composed of higher-density waters such as Bering Sea waters and winter waters (Weingartner et al., 2013), a pattern also observed during late summer in warm years (Goldstein et al., 2023). A depth of 20 m was selected since polar cod larvae primarily reside in surface waters (Geoffroy et al., 2023).

and because this depth approximates conditions in the surface mixed layer, where sea-ice melt water and ice-associated taxa might reside. Age is fish daily age. $Growth_{early}$ is otolith radius (μm) from the first 10 days of life and was included to address potential carryover effects of early life growth rates on recent growth. $PC1_{diet}$, $PC2_{diet}$, and $PC3_{diet}$ are PCA axis 1, 2, and 3 scores from the diet PCA analysis and represent a measure of diet resemblance among specimens. Variables included in the full models are those with VIFs < 3 , resulting in the exclusion of T_{200} from Equation 1 due to collinearity with T_{20} . We elected to include T_{20} as a variable in the model rather than T_{200} since it was retained in the recent growth model using Equation 2, facilitating comparisons among models with and without diet data (see Table 1 and Results).

We also tested several models that addressed potential year effects to confirm that associations between growth, environmental variability, and diet were not linked with other processes that varied among years. (1) We tested the selected model from Equation 1 excluding the year 2019 since diet differed in that year (see Results). (2) We tested the selected model from Equation 2 using only the subset of samples with diet data to facilitate more direct comparisons among models with and without diet information. (3) We compared models (Equation 1 and Equation 2) that included year as a fixed effect as well as a simple random effect (intercept; Pedersen et al., 2019) since other processes such as maternal effects, selective predation, or competition could vary by year and affect growth (Berkeley et al., 2004; Fortier and Harris, 1989; Sogard, 1997). Three years of data are fewer than the recommended group levels (5-6) for random effects (Bolker et al., 2009), but evidence suggests that there is little risk to testing models with fewer than five groups (Gelman and Hill, 2007; Gomes 2002; Oberpriller et al., 2022). For models that excluded diet data (Equation 2), the random or fixed effect of year could encompass interannual variability in diet as well as other processes (e.g. diet is an “unmeasured” variable in these models). Therefore, if an effect of year improved models that lacked diet data, then we conducted model selection on a set of models with and without a year effect to better disentangle the influence of “year” compared to temperature, salinity, age, and early growth.

GAMs were fit using restricted maximum likelihood estimation and models were compared using Akaike Information Criterion adjusted for small sample sizes (AICc; Akaike, 1998). If AICc differed by ≤ 2 then the more parsimonious model was selected. For each model

formulation, a stepwise variable selection procedure was conducted and final models were selected among the most supported candidate models (Burnham and Anderson, 2002; Zuur et al., 2009). For all models, smooth terms were replaced with a linear relationship if the estimated degrees of freedom (edf) were equal to 1. All data analysis was conducted in R (R Core Team, 2024) using packages FSA for age analyses (Ogle et al., 2025), Vegan (Oksanen et al., 2025) and MASS (Venables and Ripley, 2002) for multivariate analyses, and mgcv for GAMs (Wood, 2017).

3. Results

Spatial extent and coverage of surveys differed among the study years, but polar cod catch locations in late summer were primarily in the northeast region of the Chukchi Sea in cooler water temperatures (Fig. 1, Supplemental Table 1, Supplemental Fig. 1). Hatch dates were similar in 2017 and 2019, and later in 2018 when temperatures were also generally cooler at catch locations (Figs. 1, 2A). Fish size-at-age overlapped among years, but fish were slightly larger in 2019 (Fig. 2B). Early growth [Fig. 2C; ANOVA: $F(2, 69) = 4.99$, $p < 0.01$; Tukey: 2017 = 2018, 2018 = 2019, 2017 < 2019], past growth [Fig. 2D; ANOVA: $F(2, 67) = 20.00$, $p < 0.001$; Tukey: 2017 = 2018, 2017 & 2018 < 2019], and recent growth [Fig. 2D; ANOVA: $F(2, 69) = 9.67$, $p < 0.001$; Tukey: 2017 = 2018, 2017 & 2018 < 2019] were greatest in 2019. Trends in growth rates indicated elevated past growth compared to recent growth in 2019 (Fig. 2D). Growth rates were relatively similar across overlapping ages in 2017 and 2018 (Fig. 2E). Fish in 2019 consistently grew faster, though there was a narrower range in overlapping ages between 2019 and the other years (Fig. 2E). There were no clear spatial patterns in recent growth (Fig. 1, Supplemental Fig. 2).

Polar cod diets in 2019 were distinct from the other years (Fig. 3) as confirmed by significant differences between PC axis 1 scores [ANOVA: $F(2, 48) = 13.5$, $p < 0.001$; Tukey: 2017 = 2018, 2017 & 2018 < 2019]. This overarching variation in diet was likely minimally related to body size (diet ontogeny) or *in situ* water temperature (Fig. 3). Positive scores along PCA axis 1, where all 2019 polar cod diets ordinated, were primarily associated with the presence of the large copepod *C. glacialis* and the smaller copepods, *Oithona* spp. and *Pseudocalanus* spp. as well the absence of unidentifiable copepods (C_UNID). There was divergence along PCA axis 2 among the 2019 fish with a separation between those that primarily

consumed *C. glacialis*, and those that consumed smaller copepod taxa. PCA axis 3 captured similar patterns to PCA axis 1, but with negative scores reflecting consumption of small calanoid copepods (C_Cal_sm) and positive scores reflecting greater consumption of *C. glacialis*, *Oithona* spp., and unidentifiable copepods (Fig. 3D). PCA axis 4 explained a small proportion of variability in the data (7.72%) but primarily separated consumption of the broad taxonomic grouping of large calanoid copepods from *C. glacialis*, *Pseudocalanus* spp., and unidentified copepods. Individual fish diets varied to a greater degree in 2017 and 2018 than 2019, but some fish diets resembled those from 2019 (Fig. 3A, C).

Diet resources determined from CSIA-AA were similar in 2017 and 2018 (Fig. 4A). CSIA-AA data were not available for 2019, potentially affecting observed variation in diet resources and resulting in weak relationships between CSIA-AA and gut contents (Fig. 4B). Only a small proportion of variation in diet data was correlated with changes in CSIA-AA (32.3% based on both constrained axes); nevertheless, there were indications of potential links between diet and basal resources. CSIA-AA values for polar cod primarily overlapped with microalgae and pPOM (Fig. 4A). However, positive PCA axis 1 scores were associated with *Pseudocalanus* spp., which can include temperate and sea-ice associated species in the Chukchi Sea, as well as *C. glacialis* which is a cold-water, ice-associated copepod (Questel et al. 2016; Spear et al. 2020). These positive PCA axis 1 scores were strongly correlated with the amino acid Isoleucine and had a weak negative association with Leucine (Fig. 4). This separation between Isoleucine and Leucine was a major discriminating pattern in the CSIA-AA “fingerprint” for iPOM, which is indicative of carbon resources originating from sea-ice algae, to a greater degree than pPOM which is representative of pelagic primary producers (Fig. 4A).

Recent growth was primarily associated with variation in diet, with only weak relationships with temperature, salinity, or early growth based on comparisons of GAMs that included or excluded diet information (Table 1; Supplemental Table 3). As expected, given few group levels, models with year as a fixed or random effect showed similar results (Supplemental Tables 3 and 4; Gelman and Hill, 2007). There was little support for the inclusion of year as a random or fixed effect in the recent growth diet models (Supplemental Table 3), suggesting that accounting for unmeasured variation among years did not improve model fits. Models without diet information improved when a year effect was included, but all model formulations

corroborated the minimal influences of temperature, salinity, and early growth on recent growth (Table 1; Supplemental Table 3). The substantial impact of diet on recent growth was confirmed by several model comparisons. (1) Since diet information was not available for all samples (Supplemental Table 1), we tested “no diet” models using the identical subset of samples used in “diet” models (n=51) to facilitate comparisons between “diet” and “no diet” model results (Table 1 and Supplemental Table 3). This more direct comparison confirmed that models that included diet information showed considerable improvements over those that excluded diet data. This was especially pronounced when year was not included in the “no diet” model (Table 1), but continued to hold true when “no diet” models included a fixed or random effect of year (Supplemental Table 3; for example see Diet (Eq 1) and No diet (Eq 2) + Year_{re}: diet fish only). Since diet differed among years (e.g. 2019), this comparison also suggests that diet affected recent growth to a greater degree, or in addition to, other factors that varied among years. (2) A model that excluded data from 2019 continued to show a substantial impact of diet on recent growth. This ensured that variation in diet in 2019 compared to the other years was not driving the correlations between diet and growth (Table 1).

Recent growth had little to no relationship with age such that age was not included in the final GAMs (Table 1, Supplemental Table 3). Neither temperature nor salinity were retained in the selected recent growth diet model, nor the “no diet” models that included fixed or random effects of year (Table 1; Supplemental Table 3). Early growth was retained in the selected models that excluded diet data, but the relationship was weakly negative, suggesting that early larval growth may have a nominal carryover effect on later growth rates (Supplemental Table 3). Positive PCA axis 1 (primarily greater consumption of *C. glacialis*, *Pseudocalanus* spp., and of *Oithona* spp. and reduced consumption of unidentified copepods) and PCA axis 3 scores (also associated with greater consumption of *C. glacialis* and *Oithona* spp., as well as reduced consumption of small calanoid copepods) were associated with faster recent growth (Table 1; Fig. 5A, B). Relationships between recent growth and hatch dates within each year did not show shifts in growth patterns over the capture season or across ages as would be expected if variation in growth was due to selective mortality (Supplemental Figs. 2 and 3).

Inconsistent relationships between past and recent growth among years, including a relatively flat relationship in 2019, support the supposition that recent growth was primarily

linked with recent feeding rather than a carryover effect of prior condition and growth (Supplemental Fig. 4). Past growth was minimally correlated with temperature and salinity (Supplemental Table 4). Unlike recent growth, all past growth models that included a year effect outperformed those that did not account for year, regardless of whether diet information was incorporated in the models (Supplemental Table 4). The strong relationship between sampling year and past growth suggests that other processes that varied among years had a considerable impact on past growth. These results indicate that diet at capture explained appreciably less variation in past growth than recent growth, in accordance with the expectation that gut contents reflect recent conditions since evacuation occurs on the order of hours to days depending on feeding, temperature, and development (Canino and Bailey, 1995; Hop and Tonn, 1998; Table 1, Supplemental Tables 3 and 4).

4. Discussion

The relative influences of temperature and diet on growth of marine fish is a fundamental ecological question that continues to gain importance in polar regions as sea ice declines and ocean temperatures increase, potentially leading to shifts in phenology, growth rates, and predator-prey mismatch (Atkinson, 1994; David et al., 2022; Geoffroy et al., 2023; Mazur et al., 2007; Meekan et al., 2003). A warming Arctic will likely exacerbate metabolic constraints on growth of cold-adapted species through the combined effects of thermal tolerances and food quality and availability. The relationship between polar cod growth and temperature was weak despite evidence of thermal optima in laboratory experiments (Koenker et al., 2018; Laurel et al., 2016), the potential for interactions between temperature and prey quality, and expectations that reduced sea-ice cover and warming temperatures may limit access to favorable environments (Geoffroy et al., 2023). While lagged effects on growth could differ between temperature and diet, our results suggest that diet had a greater influence on larval and juvenile growth rates than temperature during the three warm study years, largely driven by the consumption of lipid-rich copepods such as *C. glacialis*. These findings suggest that in a warm Arctic, the presence of high-quality prey can support fast growth of larval and juvenile polar cod and buffer against potential negative effects of sea-ice decline.

Variation in diet and growth within and among years suggests that access to habitat patches with cooler temperatures and high-quality prey is essential to the adaptability of polar cod to a warming climate. Even during warm years, polar cod were primarily collected in colder water temperatures and at high latitudes (e.g., ~71°N or higher), suggesting differentiation in habitat use that has also been observed for older age-0 fish during colder years in the Chukchi Sea (Marsh et al., 2020). While 2019 was ostensibly the warmest year of the study (Stabeno et al., 2023), polar cod had faster growth and consumed higher quality prey species that are typically associated with Arctic water masses and sea ice. This dichotomy suggests that fish were accessing, or co-occurring with, patches of favorable prey in 2019 despite warm temperatures. During 2017 and 2018, there was greater variation in diet among individuals compared to 2019, yet fish in favorable habitats with high-quality prey grew faster regardless of the year. This pattern suggests that overall slower growth in 2017 and 2018 was driven by variation in feeding habitat, whereby a greater majority of the sampled fish encountered poorer habitats in 2017 and 2018 than in 2019. Juvenile polar cod also had low body condition in 2017, but high body condition in 2019 that was comparable to favorable cold years and attributed to localized conditions (Copeman et al., 2022a). These contrasting warm years suggest that annual variation in climate may be mediated by localized processes and highlight the importance of remaining, or remnant, sea-ice habitats and associated prey to support polar cod growth and survival in a “future” Arctic.

CSIA-AA results supported the link between favorable diets and lingering sea-ice productivity to support fast growth. CSIA-AA $\delta^{13}\text{C}$ from fish tissue reflects basal carbon resources that are influenced by prey consumption and interim trophic linkages. For example, polar cod prey such as *C. glacialis* utilizes a substantial portion of carbon from sea-ice environments (Kohlbach et al., 2016). However, they consume a variety of prey such as microzooplankton, phytoplankton, and sea-ice algae (Campbell et al., 2009; Sherr et al., 2009; Søreide et al., 2010), and reliance on sympagic carbon resources can shift seasonally for *Calanus* species in Arctic food webs (Kohlbach et al., 2024). In warming temperatures, polar cod and their prey may both exhibit diet shifts away from sea-ice resources. Nonetheless, correlations between favorable diets and sea-ice carbon resources were positive, despite lacking data from 2019 when diets diverged from other years and relationships between diet and growth were most prominent. This evidence for lingering sea-ice basal carbon resources in polar cod diets further

supports the important role of sea-ice productivity across trophic levels and alludes to the continuing importance of sea-ice carbon resources in Arctic food webs during late summer in warm years.

This study was limited to only warm years in the Chukchi Sea and relatively small sample sizes, necessitating future efforts to contrast the influences of diet and temperature on larval and juvenile polar cod growth across broader environmental and geographic domains. However, the Chukchi Sea ecosystem represents a productive shallow shelf environment that functions as a transitional zone between subarctic and Arctic habitats. As a result, this region is changing at a rapid pace and may provide insight into potential consequences of continuing Arctic sea-ice recession. Comparisons among the study years indicate conditions that promote growth and survival of polar cod in a warming Arctic. For example, reduced sea-ice cover combined with wind reversals in 2017 (Goldstein et al., 2023) potentially impacted distributions of polar cod and their prey, and limited access to ice-associated habitats. This contrasts with 2019 when winds were more consistent, and water masses and temperature showed stronger latitudinal gradients (Levine et al., 2023), potentially supporting remnant sea-ice habitats at higher latitudes despite low ice extent. Notably, variation in diet was strongly correlated with recent growth but less tightly coupled to growth beyond 5-d prior to capture, suggesting that encounters with suitable habitat and high-quality prey could have an expeditious effect on larval and juvenile polar cod growth. The capacity for polar cod to exploit available favorable environments in exceptionally warm years points to adaptability and resilience of the species, but suggests that loss of sea-ice habitats earlier in the warm season and diminished access to high-quality prey have detrimental effects on growth of an important Arctic forage fish species.

Data statement

Data will be made available on request.

Acknowledgments

We are grateful for the expertise and insightful reviews of this manuscript provided by Robert Levine and Cynthia Yeung in addition to the thoughtful comments of four anonymous reviewers. We thank the Captains and crews of the RV *Ocean Starr* and the USCGC *Healy*. We also thank

NOAA's Alaska Fisheries Science Center's Ecosystems and Fisheries-Oceanography Coordinated Investigations Program (EcoFOCI) and Midwater Assessment and Conservation Engineering Program for their efforts at sea and their taxonomic expertise. This manuscript is EcoFOCI Contribution EcoFOCI-XXXX. The findings and conclusions in the paper are those of the authors and do not necessarily represent the views of the National Marine Fisheries Service, NOAA. Reference to trade names does not imply endorsement by the National Marine Fisheries Service, NOAA.

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E.D. Goldstein: Conceptualization, Data curation, Formal analysis, Investigation, Methodology, Writing – original draft, Writing – review and editing. K.E. Axler: Conceptualization, Data curation, Investigation, Methodology, Writing – review and editing. J.F. Lamb: Data curation, Investigation, Methodology, Writing – review and editing. M.C. Rogers: Data curation, Investigation, Methodology, Writing – review and editing. J.T. Duffy-Anderson: Conceptualization, Investigation, Funding acquisition, Methodology, Resources, Writing – review and editing

Funding sources

This research was supported by the North Pacific Research Board's Arctic Integrated Ecosystem Research Program (AIERP; <https://www.nprb.org/arctic-program/>). Additional funding was received from the Arctic Research Program, NOAA Research.

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Tables

Table 1. Generalized additive models (GAMs) linking recent growth (average otolith increment width for the most recent 5 days prior to capture, $\mu\text{m}/\text{day}$) with intrinsic factors, environmental factors, and diet from gut content analysis (diet). Model description identifies whether the full model included diet information, references the model equation [Equation 1 (Eq 1) or Equation 2 Eq 2)], and provides additional model descriptions. Rank identifies the top two models (selected model in bold) for each model description category. Two alternative (alt) models are also presented to confirm that patterns were consistent among relevant subsets of the data: fish from 2017 and 2018 only (no 2019) and diet fish only. Model lists the smooth terms [s()] and parametric terms in each model. PC1_{diet} and PC3_{diet} are PCA axis 1 and 3 scores from the diet PCA analysis (see Fig. 3). $\text{Growth}_{\text{early}}$ is otolith radius from the first 10 days of life. T_{20} and S_{200} are average temperature from the surface to a depth of 20 m and average salinity to a depth of 200 m, respectively. Adjusted coefficient of determination (R^2), Akaike information criterion adjusted for small sample sizes (AICc), and sample size (N) are shown for all models. AIC is not provided for the Diet (Eq 1): no 2019 model since the sample subset differs from all other models presented. See Supplemental Table 3 for additional model comparisons.

Model description	Rank	Model	R^2	AICc	N
Diet (Eq 1)	1	s(PC1_{diet}) + s(PC3_{diet})	0.43	35.81	51
	2	s(PC1 _{diet}) + s(PC3 _{diet}) + $\text{Growth}_{\text{early}}$	0.45	35.10	51
No diet (Eq 2)	1	T_{20}	0.06	69.57	72
	2	$S_{200} + T_{20}$	0.07	70.12	72
Diet (Eq 1): no 2019	alt	s(PC1 _{diet}) + s(PC3 _{diet})	0.30	-	41
No diet (Eq 2): diet fish only	alt	T_{20}	0.13	52.61	51

Figure Captions

Figure 1. Polar cod (*Boreogadus saida*) catch locations in the Chukchi Sea from three years (2017-2019; circle symbols) and interpolated co-collected CTD temperature data from the same surveys averaged over the top 20 m of the water column (“x” symbols show station locations) to correspond with generalized additive model results (Table 1; see Supplemental Fig. 1 for average temperature over the top 200 m). The color scale for fish catch locations shows average otolith-derived growth ($\mu\text{m}/\text{day}$) for the most recent 5 days of life preceding capture. The size of circles indicates otolith-derived age of each sample and circles are jittered to minimize overlap.

732

733 Figure 2. Otolith-derived hatch and growth information for polar cod by year. (A) Kernel density
 734 distribution of hatch day of year where n is sample size (see Supplemental Table 1). (B) Age
 735 versus length where line fits are a Gompertz function (solid; root mean square error=3.79; Ogle
 736 et al., 2025) and a linear fit (dotted; root mean square error=3.94). (C) Otolith radius from the
 737 otolith core through the first 10 days of life (early growth). Boxplots show the median (line), the
 738 1st and 3rd quartiles (hinges), and upper and lower whiskers that extend from the hinge to the
 739 largest (upper) and smallest (lower) value no further than 1.5 times the inter-quartile range, and
 740 outliers beyond the whiskers (points). (D) Mean growth ($\pm$ SD) measured from the first full day
 741 prior to capture (day 1) until increments were no longer interpretable at the otolith edge focal
 742 plane. The dark gray line delineates the first 10 days prior to capture. The light gray line (15
 743 days) shows the 1st quartile of increments measured from the edge. (E) Mean otolith increment
 744 width ($\pm$ SD) by back-calculated age. Age was back-calculated using catch date, fish age, and
 745 increment count from the otolith edge. Back-calculated ages are averaged for ages with at least 3
 746 observations and exclude increment widths greater than 15 days prior to capture (e.g., 1st
 747 quartile) since few specimens had measurements beyond that window of time.

748

749 Figure 3. Principal component (PC) analysis of gut content diet data from larval and juvenile
 750 polar cod (*Boreogadus saida*) showing PC axes 1 and 2 (A, B) and PC axes 3 and 4 (C, D). (A)
 751 and (C) show specimen scores (scaled to preserve Euclidean distance between points) color
 752 coded by year. (B) and (D) show vector loadings (bold arrows) with labels that describe the
 753 copepod prey taxa (scaled to ensure that angles between vectors reflect their correlations). To aid
 754 in interpretation, dotted line segments connect offset labels with corresponding vectors loadings.
 755 Symbol size in (A) and (C) corresponds to fish standard length (mm). The color scale in (B) and
 756 (D) shows interpolated temperature data averaged throughout the water column from each catch
 757 location to visualize temperature associations of prey items (interpolations correspond to points
 758 in A and C but are scaled to correspond with B and D). Percentages refer to the amount of
 759 variability in the data described by each PC axis. Abbreviated taxa include taxonomic
 760 descriptions where the prefix “C_” designates copepods. Abbreviations are provided in
 761 Supplemental Table 2.

Figure 4. (A) Linear Discriminant Analysis (LDA), adapted from Goldstein et al. (2023) to visualize relationships between compound-specific stable isotopes of amino acids (CSIA-AA $\delta^{13}\text{C}$) from polar cod tissue, sea-ice particulate organic matter (iPOM), pelagic particulate organic matter (pPOM), and potential carbon sources used to develop the LDA [Microalgae, *Synechococcus* spp. (Syn) combined with Dinoflagellates (Dino), and bacteria]. CSIA-AA $\delta^{13}\text{C}$ from polar cod, iPOM, and pPOM are predictions (solid symbols) and polar cod specimens (labeled 2017 and 2018) are color coded by collection year. (B) Redundancy analysis triplot showing the influence of polar cod diet composition (PC scores 1-3 shown in Fig. 3) on CSIA-AA $\delta^{13}\text{C}$. Points are polar cod specimens and symbol size corresponds with fish age. Black vectors are explanatory variables. Gray vectors are response variables where angles between vectors approximate their linear correlations. In (A) and (B) essential amino acids from CSIA-AA $\delta^{13}\text{C}$ used for analyses are abbreviated as Val (Valine), Leu (Leucine), and Ile (Isoleucine).

Figure 5. Generalized additive model partial effects plots [Table 1: Diet (Eq 1) selected model] showing relationships between otolith-derived recent growth ($\mu\text{m}/\text{day}$; increment widths averaged over 5 days prior to capture) and A) PC1 diet scores and B) PC3 diet scores. Smooth terms are denoted by s() and gray bands are 2 standard errors above and below the estimates of the terms.

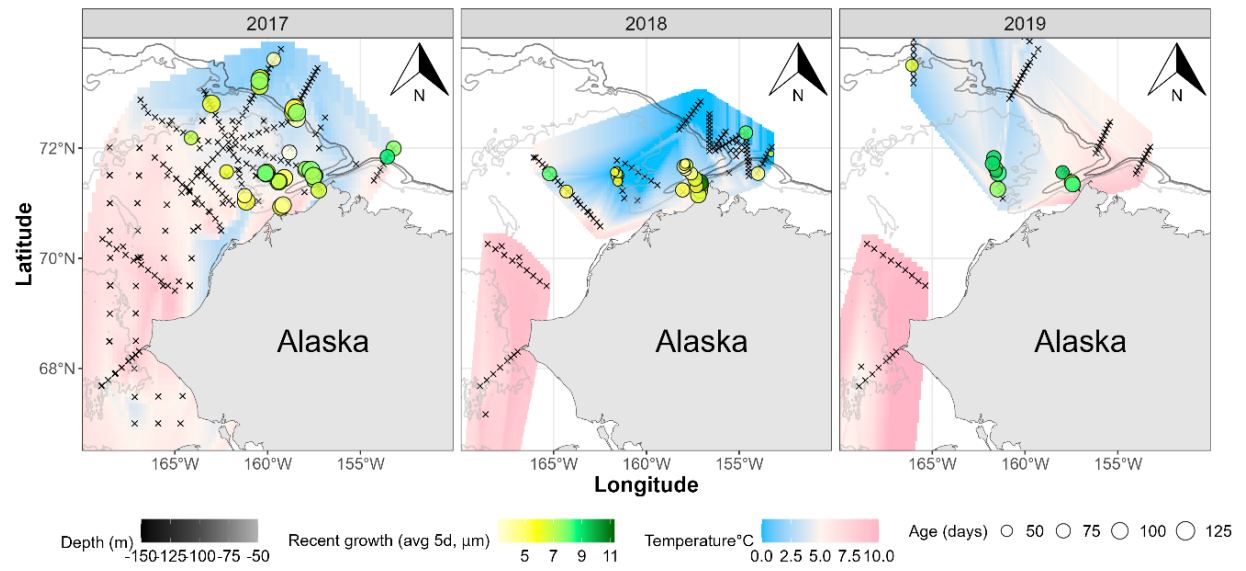


Figure 1.

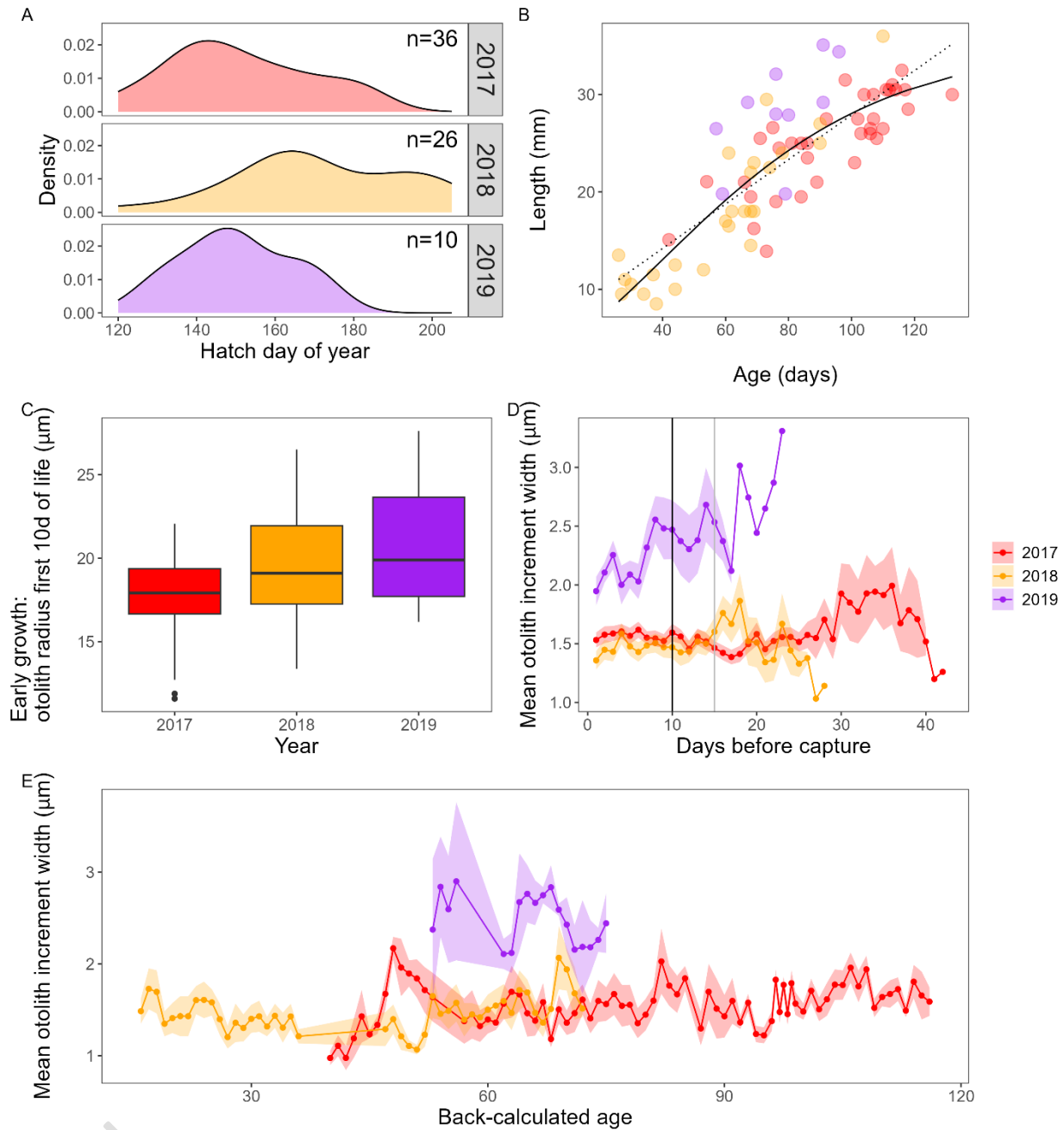


Figure 2.

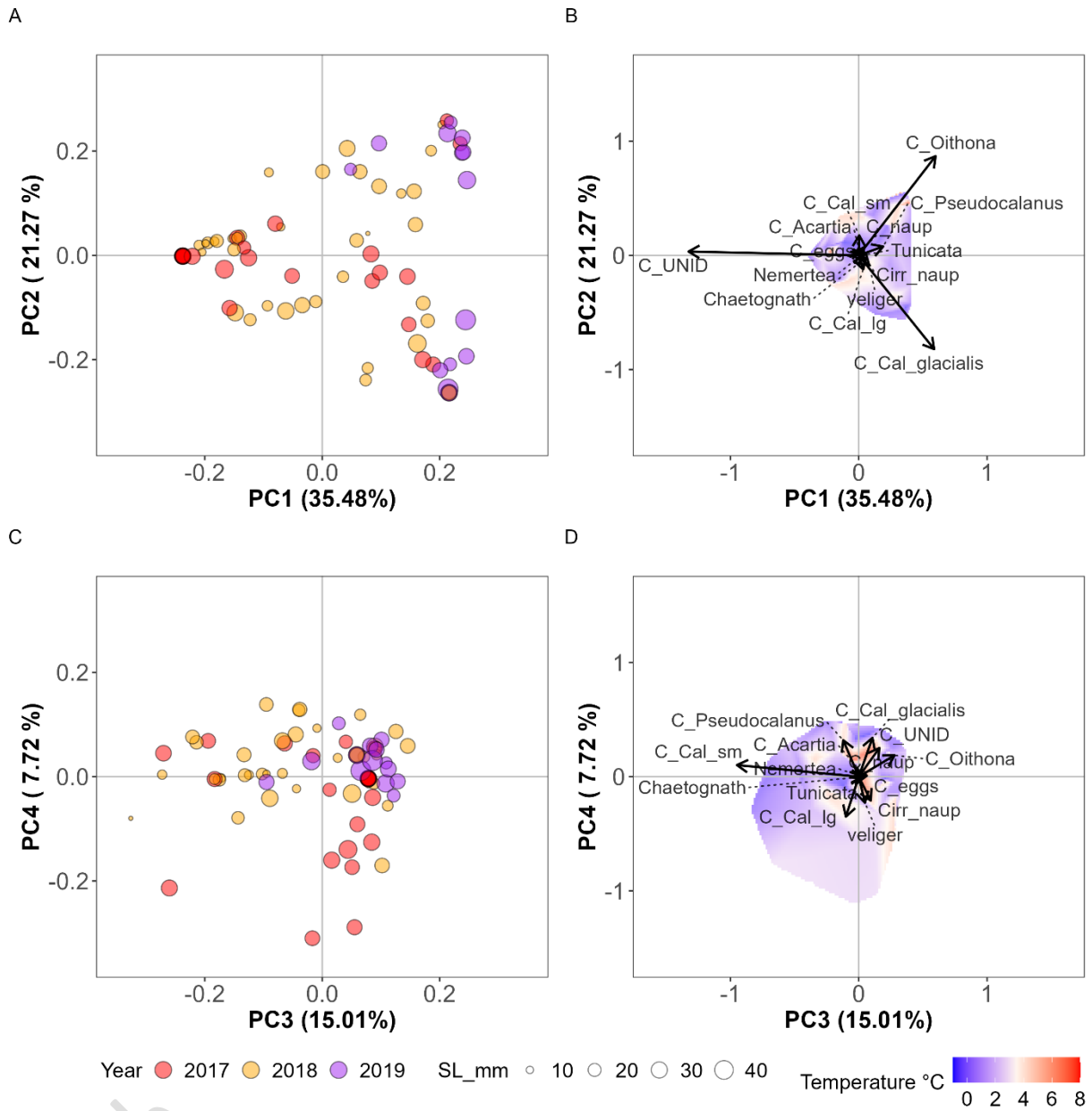


Figure 3.

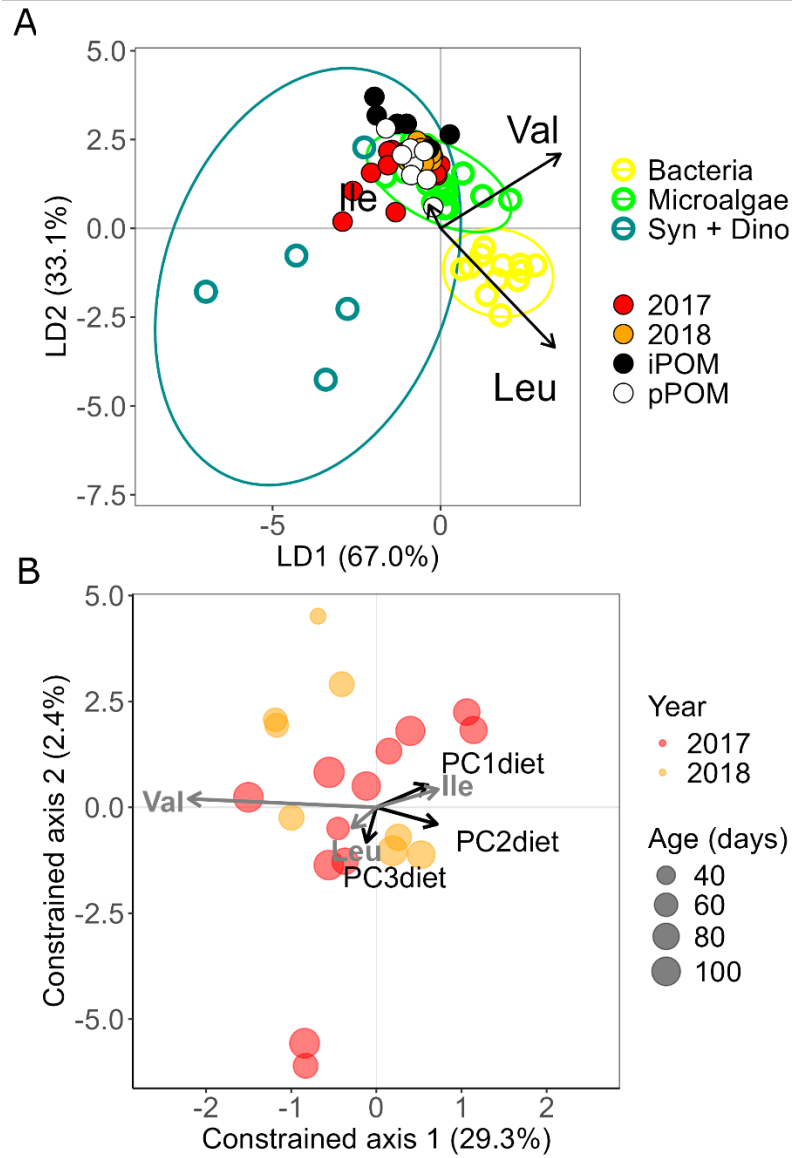
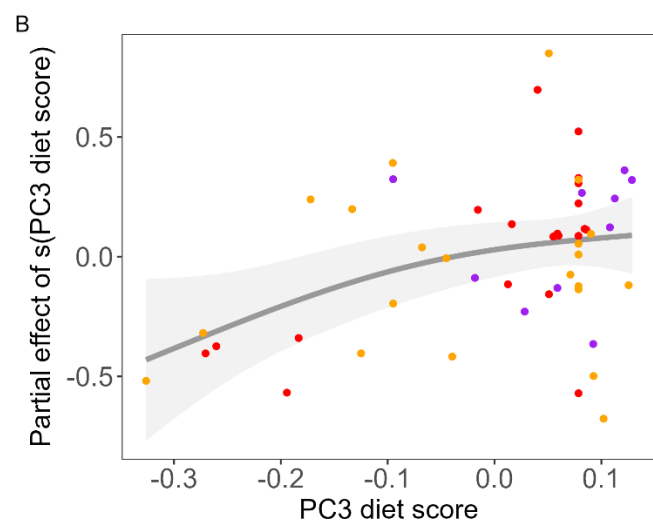
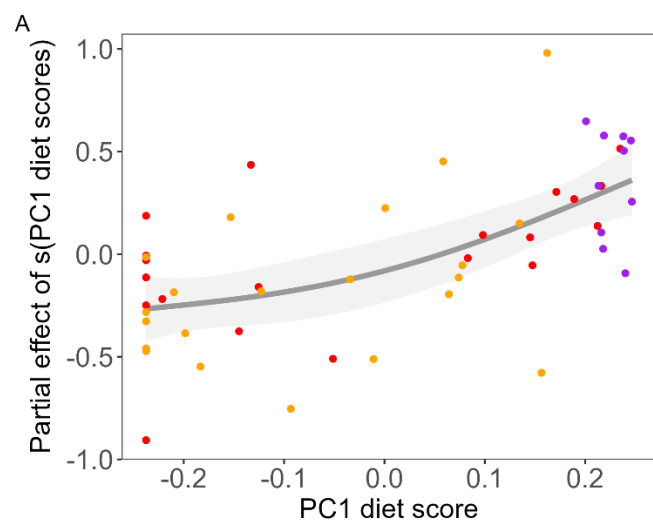


Figure 4.



• 2017 • 2018 • 2019

Figure 5.